





Nucleotide

Protein

4788 bp

Genome Structure PopSet

Taxonomy

OMIM

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08-NOV-1994

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orthographic and the second

Human glucocorticoid □ 1: M10901 receptor alpha mRNA,

PubMed, Protein, Related Sequences, Taxonomy, OMIM, LinkOut

complete cds

HUMGCRA

DEFINITION ACCESSION

Human glucocorticoid receptor alpha mRNA, complete cds. M10901

VERSION

M10901.1 GI:183032

KEYWORDS

glucocorticoid receptor; glucocorticoid receptor-alpha.

mRNA

SOURCE

LOCUS

Human lymphoid cell line IM-9, cDNA to mRNA, clones hGR[1.2, 2.9, 5.16] and fibroblast cDNA library (H.Okayama), clones OB7 and OB10.

ORGANISM

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE

(bases 1 to 4788)

AUTHORS

TITLE

Hollenberg, S.M., Weinberger, C., Ong, E.S., Cerelli, G., Oro, A.,

Lebo, R., Thompson, E.B., Rosenfeld, M.G. and Evans, R.M. Primary structure and expression of a functional human

glucocorticoid receptor cDNA

JOURNAL

Nature 318 (6047), 635-641 (1985)

MEDLINE

86092206

COMMENT

Although [1] did not actually sequence this entire sequence from one contiguous clone (thus eliminating the possibility that this sequence contains segments from multiple genes) their evidence strongly suggests that the alpha clone, OB7, and the beta clone, OB10, are transcribed from the same gene on chromosome 5 by alternate mRNA splicing. Therefore, the predicted alpha mRNA sequence is presented here in its entirety. Positions 1 to 648 were determined from clones OB10 and hGR5.16; positions 649-4788 were determined by overlapping regions of all 5 clones listed on the SOURCE line. The beta clone diverges from this alpha clone in sequence after position 2313. See also the beta GCR mRNA in entry with accession number M11050. [1] reports that the alpha form of glucocorticoid receptor is the predominant physiological form found in the various human and mouse cell lines that they tested. [1] also noted a region of chromosome 16 with enough homology to these clones to hybridize efficiently. Alternate polyadenylation signals present at positions 3101-3106 and 4678-4684 may also be utilized by some mRNAs. A clone OB12 was isolated that used the 3101-3106 signal.

FEATURES

Location/Qualifiers

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/map = "5q31 - q32"

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CDS

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http://www.../query.fcgi?cmd=Retrieve&db=Nucleotide&list_uids=183032&dopt=GenBan \ 02/15/2001

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BASE COUNT
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Nucleotide Protein Genome Structure **PopSet** Taxonomy **OMIM** Search Nucleotide Limits History Index Clipboard Default View as HTML (SE)1/E arcelevie course at le Human mRNA for beta-PubMed, Protein, Related Sequences, Taxonomy, OMIM, LinkOut glucocorticoid receptor (clone **OB10**) LOCUS **HSGCRBR** 3791 bp mRNA PRI 12-SEP-1993 DEFINITION Human mRNA for beta-glucocorticoid receptor (clone OB10). ACCESSION X03348 M11050 VERSION X03348.1 GI:31681 KEYWORDS glucocorticoid receptor. SOURCE human. ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. REFERENCE (bases 1 to 3791) Hollenberg, S.M., Weinberger, C., Ong, E.S., Cerelli, G., Oro, A., **AUTHORS** Lebo, R., Thompson, E.B., Rosenfeld, M.G. and Evans, R.M. TITLE Primary structure and expression of a functional human glucocorticoid receptor cDNA JOURNAL Nature 318 (6047), 635-641 (1985) MEDLINE 86092206 **FEATURES** Location/Qualifiers source 1..3791 /organism="Homo sapiens" /db xref="taxon:9606" CDS 133..2361 /note="(aa 1-742)" /codon start=1 /product="beta-glucocorticoid receptor" /protein id="CAA27054.1" /db xref="GI:31682" /db xref="SWISS-PROT:P04150" /translation="MDSKESLTPGREENPSSVLAQERGDVMDFYKTLRGGATVKVSAS SPSLAVASQSDSKQRRLLVDFPKGSVSNAQQPDLSKAVSLSMGLYMGETETKVMGNDL GFPQQGQISLSSGETDLKLLEESIANLNRSTSVPENPKSSASTAVSAAPTEKEFPKTH SDVSSEQQHLKGQTGTNGGNVKLYTTDQSTFDILQDLEFSSGSPGKETNESPWRSDLL IDENCLLSPLAGEDDSFLLEGNSNEDCKPLILPDTKPKIKDNGDLVLSSPSNVTLPQV KTEKEDF1ELCTPGV1KQEKLGTVYCQASFPGAN1IGNKMSA1SVHGVSTSGGQMYHY DMNTASLSQQQDQKPIFNVIPPIPVGSENWNRCQGSGDDNLTSLGTLNFPGRTVFSNG YSSPSMRPDVSSPPSSSSTATTGPPPKLCLVCSDEASGCHYGVLTCGSCKVFFKRAVE GQHNYLCAGRNDCIIDKIRRKNCPACRYRKCLQAGMNLEARKTKKKIKGIQQATTGVS QETSENPGNKTIVPATLPQLTPTLVSLLEVIEPEVLYAGYDSSVPDSTWRIMTTLNML GGRQVIAAVKWAKAIPGFRNLHLDDQMTLLQYSWMFLMAFALGWRSYRQSSANLLCFA PDLIINEQRMTLPCMYDQCKHMLYVSSELHRLQVSYEEYLCMKTLLLLSSVPKDGLKS QELFDEIRMTYIKELGKAIVKREGNSSQNWQRFYQLTKLLDSMHENVMWLKPESTSHT LI" <u>misc</u> feature 3168..3173 /note="pot. polyA signal" 3539..3545 <u>misc</u> feature /note="pot. polyA signal" misc_feature 3770..3775 /note="put. polyA signal" 3791 poly<u>A</u> site 759 c BASE COUNT 1162 a 808 g 1062 t ORIGIN 1 tttttagaaa aaaaaaatat atttccctcc tgctccttct gcgttcacaa gctaagttgt

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